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Gln Glu Leu Ser Leu Arg Leu Lys Ser Cys Phe Thr Lys Asp Tyr Glu 115 120 125

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ctc tat gtc aaa gac cct gcc cgg ccc tgg aac gtg cta gca cag gag Leu Tyr Val Lys Asp Pro Ala Arg Pro Trp Asn Val Leu Ala Gln Glu 100 105 110	634
gtg gtc gtg ttc gag gac cag gac gca cta ctg ccc tgt ctg ctc aca Val Val Phe Glu Asp Gln Asp Ala Leu Leu Pro Cys Leu Leu Thr 115 120 125 130	682

															cgg Arg	730
	ctc Leu	_	_							_						778
	atc Ile															826
	ctg Leu 180															874
	cag Gln															922
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	agc Ser															1018
aag Lys	ctc Leu	gca Ala 245	atc Ile	cct Pro	caa Gln	·caa Gln	tct Ser 250	gac Asp	ttt Phe	cat His	aat Asn	aac Asn 255	cgt Arg	tac Tyr	caa Gln	1066
aaa Lys	gtc Val 260	ctg Leu	acc Thr	ctc Leu	aac Asn	ctc Leu 265	gat Asp	caa Gln	gta Val	gat Asp	ttc Phe 270	caa Gln	cat His	gcc Ala	ggc Gly	1114
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atg Met	ttc Phe	ttc Phe	cgg Arg	gtg Val 295	gta Val	gag Glu	agt Ser	gcc Ala	tac Tyr 300	ttg Leu	aac Asn	ttg Leu	agc Ser	tct Ser 305	gag Glu	1210
cag Gln	aac Asn	ctc Leu	atc Ile 310	cag Gln	gag Glu	gtg Val	acc Thr	gtg Val 315	gly aaa	gag Glu	gjå aaa	ctc Leu	aac Asn 320	ctc Leu	aaa Lys	1258
gtc Val	atg Met	gtg Val 325	gag Glu	gcc Ala	tac Tyr	cca Pro	ggc 330	ctg Leu	caa Gln	ggt Gly	ttt Phe	aac Asn 335	tgg Trp	acc Thr	tac Tyr	1306
ctg Leu	gga Gly 340	ccc Pro	ttt Phe	tct Ser	gac Asp	cac His 345	cag Gln	cct Pro	gag Glu	ccc Pro	aag Lys 350	ctt Leu	gct Ala	aat Asn	gct Ala	1354
	acc Thr															1402
ctg Leu	aag Lys	ccc Pro	tct Ser	gag Glu	gct Ala	ggc Gly	cgc Arg	tac Tyr	tcc Ser	ttc Phe	ctg Leu	gcc Ala	aga Arg	aac Asn	cca Pro	1450

				375					380					385			
															cca Pro	149	98
	_	_	_												ttg Leu	154	16
	gct Ala 420															159	94
	ggc Gly															164	12
	gac Asp															169	90
	gtg Val															173	88
tac Tyr	gag Glu	tgc Cys 485	agg Arg	gcc Ala	cac His	aac Asn	agc Ser 490	gtg Val	Gly ggg	agt Ser	ggc Gly	tcc Ser 495	tgg Trp	gcc Ala	ttc Phe	178	36
ata Ile	ccc Pro 500	atc Ile	tct Ser	gca Ala	gga Gly	gcc Ala 505	cac His	acg Thr	cat His	ccc Pro	ccg Pro 510	gat Asp	gag Glu	ttc Phe	ctc Leu	183	34
ttc Phe 515	aca Thr	cca Pro	gtg Val	gtg Val	gtc Val 520	gcc Ala	tgc Cys	atg Met	tcc Ser	atc Ile 525	atg Met	gcc Ala	ttg Leu	ctg Leu	ctg Leu 530	188	32
ctg Leu	ctg Leu	ctc Leu	ctg Leu	ctg Leu 535	cta Leu	ttg Leu	tac Tyr	aag Lys	tat Tyr 540	aag Lys	cag Gln	aag Lys	ccc Pro	aag Lys 545	tac Tyr	193	0 0
cag Gln	gtc Val	cgc Arg	Trp	Lys	atc Ile	Ile	Glu	Ser	Tyr	Glu	Gly	aac Asn	Ser	Tyr	act Thr	197	'8
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cgg Arg	aac Asn 580	aac Asn	ctg Leu	cag Gln	ttt Phe	ggt Gly 585	aag Lys	acc Thr	ctc Leu	gga Gly	gct Ala 590	gga Gly	gcc Ala	ttt Phe	Gly 333	207	4
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Lys	Glu	Ala	Leu 630	Met	Ser	Glu	Leu	Lys 635	Ile	Met	Ser	His	Leu 640	Gly	Gln	
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					gag Glu 680											2362
					Gly											2410
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					cac His											2650
					gcc Ala											2698
- 7		-	Ξ	_	aac Asn		_ ~		_	===	_	=				2746
					cca Pro											2794
					tcc Ser 840											2842
					tac Tyr											2890
					gga Gly											2938

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cac aga ccc acc ttc cag cag atc tgc tcc ttc ctt cag gag cag gcc His Arg Pro Thr Phe Gln Gln Ile Cys Ser Phe Leu Gln Glu Gln Ala 900 905 910	3034
caa gag gac agg aga gag cgg gac tat acc aat ctg ccg agc agc agc Gln Glu Asp Arg Arg Glu Arg Asp Tyr Thr Asn Leu Pro Ser Ser Ser 915 920 925 930	3082
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tct agt gag cac ctg acc tgc tgc gag caa ggg gat atc gcc cag ccc Ser Ser Glu His Leu Thr Cys Cys Glu Gln Gly Asp Ile Ala Gln Pro 950 955 960	3178
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Gly Gln Gly Ile Pro Val Ile Glu Pro Ser Val Pro Glu Leu Val Val

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Lys Pro Gly Ala Thr Val Thr Leu Arg Cys Val Gly Asn Gly Ser Val 35 40 45

Glu Trp Asp Gly Pro Pro Ser Pro His Trp Thr Leu Tyr Ser Asp Gly 50 55 60

Ser Ser Ser Ile Leu Ser Thr Asn Asn Ala Thr Phe Gln Asn Thr Gly 65 70 75 80

Thr Tyr Arg Cys Thr Glu Pro Gly Asp Pro Leu Gly Gly Ser Ala Ala 85 90 95

Ile His Leu Tyr Val Lys Asp Pro Ala Arg Pro Trp Asn Val Leu Ala
100 105 110

Gln Glu Val Val Phe Glu Asp Gln Asp Ala Leu Leu Pro Cys Leu 115 120 125

Leu Thr Asp Pro Val Leu Glu Ala Gly Val Ser Leu Val Arg Val Arg 130 135 140

Gly Arg Pro Leu Met Arg His Thr Asn Tyr Ser Phe Ser Pro Trp His 145 150 155 160

Gly Phe Thr Ile His Arg Ala Lys Phe Ile Gln Ser Gln Asp Tyr Gln
165 170 175

Cys Ser Ala Leu Met Gly Gly Arg Lys Val Met Ser Ile Ser Ile Arg 180 185 190

Leu Lys Val Gln Lys Val Ile Pro Gly Pro Pro Ala Leu Thr Leu Val 195 200 205

Pro Ala Glu Leu Val Arg Ile Arg Gly Glu Ala Ala Gln Ile Val Cys 210 215 220

Ser Ala Ser Ser Val Asp Val Asn Phe Asp Val Phe Leu Gln His Asn 225 230 235 240

Asn Thr Lys Leu Ala Ile Pro Gln Gln Ser Asp Phe His Asn Asn Arg 245 250 255

Tyr Gln Lys Val Leu Thr Leu Asn Leu Asp Gln Val Asp Phe Gln His 260 265 270

Ala Gly Asn Tyr Ser Cys Val Ala Ser Asn Val Gln Gly Lys His Ser 275 280 285

Thr Ser Met Phe Phe Arg Val Val Glu Ser Ala Tyr Leu Asn Leu Ser 290 295 300

Ser Glu Gln Asn Leu Ile Gln Glu Val Thr Val Gly Glu Gly Leu Asn 305 310 315 320

Leu Lys Val Met Val Glu Ala Tyr Pro Gly Leu Gln Gly Phe Asn Trp 325 330 335

Thr Tyr Leu Gly Pro Phe Ser Asp His Gln Pro Glu Pro Lys Leu Ala 340 345 350

Asn Ala Thr Thr Lys Asp Thr Tyr Arg His Thr Phe Thr Leu Ser Leu 355 360 365

Pro Arg Leu Lys Pro Ser Glu Ala Gly Arg Tyr Ser Phe Leu Ala Arg 370 375 380

Asn Pro Gly Gly Trp Arg Ala Leu Thr Phe Glu Leu Thr Leu Arg Tyr 385 390 395 400

Pro Pro Glu Val Ser Val Ile Trp Thr Phe Ile Asn Gly Ser Gly Thr 405 410 415

Leu Leu Cys Ala Ala Ser Gly Tyr Pro Gln Pro Asn Val Thr Trp Leu 420 425 430

Val Trp Asp Asp Pro Tyr Pro Glu Val Leu Ser Gln Glu Pro Phe His 450 455 460

Lys Val Thr Val Gln Ser Leu Leu Thr Val Glu Thr Leu Glu His Asn 465 470 475 480

Gln Thr Tyr Glu Cys Arg Ala His Asn Ser Val Gly Ser Gly Ser Trp 485 490 495

Ala Phe Ile Pro Ile Ser Ala Gly Ala His Thr His Pro Pro Asp Glu 500 505 510

Phe Leu Phe Thr Pro Val Val Val Ala Cys Met Ser Ile Met Ala Leu 515 520 525

Leu Leu Leu Leu Leu Leu Leu Tyr Lys Tyr Lys Gln Lys Pro 530 535 540

- Lys Tyr Gln Val Arg Trp Lys Ile Ile Glu Ser Tyr Glu Gly Asn Ser 545 550 555 560
- Tyr Thr Phe Ile Asp Pro Thr Gln Leu Pro Tyr Asn Glu Lys Trp Glu 565 570 575
- Phe Pro Arg Asn Asn Leu Gln Phe Gly Lys Thr Leu Gly Ala Gly Ala 580 585
- Phe Gly Lys Val Val Glu Ala Thr Ala Phe Gly Leu Gly Lys Glu Asp 595 600 605
- Ala Val Leu Lys Val Ala Val Lys Met Leu Lys Ser Thr Ala His Ala 610 615 620
- Asp Glu Lys Glu Ala Leu Met Ser Glu Leu Lys Ile Met Ser His Leu 625 630 635 640
- Gly Gln His Glu Asn Ile Val Asn Leu Leu Gly Ala Cys Thr His Gly 645 650 . 655
- Gly Pro Val Leu Val Ile Thr Glu Tyr Cys Cys Tyr Gly Asp Leu Leu 660 670
- Asn Phe Leu Arg Arg Lys Ala Glu Ala Met Leu Gly Pro Ser Leu Ser 675 680 685
- Pro Gly Gln Asp Pro Glu Gly Gly Val Asp Tyr Lys Asn Ile His Leu 690 695 700
- Glu Lys Lys Tyr Val Arg Arg Asp Ser Gly Phe Ser Ser Gln Gly Val
 705 710 715 720
- Asp Thr Tyr Val Glu Met Arg Pro Val Ser Thr Ser Ser Asn Asp Ser 725 730 735
- Phe Ser Glu Gln Asp Leu Asp Lys Glu Asp Gly Arg Pro Leu Glu Leu 740 745 750
- Arg Asp Leu Leu His Phe Ser Ser Gln Val Ala Gln Gly Met Ala Phe 755 760 765
- Leu Ala Ser Lys Asn Cys Ile His Arg Asp Val Ala Ala Arg Asn Val 770 775 780

Leu Leu Thr Asn Gly His Val Ala Lys Ile Gly Asp Phe Gly Leu Ala 785 790 795 800

- Arg Asp Ile Met Asn Asp Ser Asn Tyr Ile Val Lys Gly Asn Ala Arg 805 810 815
- Leu Pro Val Lys Trp Met Ala Pro Glu Ser Ile Phe Asp Cys Val Tyr 820 825 830
- Thr Val Gln Ser Asp Val Trp Ser Tyr Gly Ile Leu Leu Trp Glu Ile 835 840 845
- Phe Ser Leu Gly Leu Asn Pro Tyr Pro Gly Ile Leu Val Asn Ser Lys-850 855
 - Phe Tyr Lys Leu Val Lys Asp Gly Tyr Gln Met Ala Gln Pro Ala Phe 865 870 875 880
- Ala Pro Lys Asn Ile Tyr Ser Ile Met Gln Ala Cys Trp Ala Leu Glu 885 890 895
- Pro Thr His Arg Pro Thr Phe Gln Gln Ile Cys Ser Phe Leu Gln Glu
 900 905 910
- Gln Ala Gln Glu Asp Arg Arg Glu Arg Asp Tyr Thr Asn Leu Pro Ser 915 920 925
- Ser Ser Arg Ser Gly Gly Ser Gly Ser Ser Ser Glu Leu Glu Glu 930 935 940
- Glu Ser Ser Ser Glu His Leu Thr Cys Cys Glu Gln Gly Asp Ile Ala 945 950 955 960
- Gln Pro Leu Gln Pro Asn Asn Tyr Gln Phe Cys 965 970